

# Bader Almuzzaini

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7744239/publications.pdf>

Version: 2024-02-01

21  
papers

414  
citations

933447

10  
h-index

794594

19  
g-index

22  
all docs

22  
docs citations

22  
times ranked

436  
citing authors

#	ARTICLE	IF	CITATIONS
1	Epidemiology of cancer in Saudi Arabia thru 2010â€“2019: a systematic review with constrained meta-analysis&lt;span style="font-size: 12px"&gt;& Running title: epidemiology of cancer in Saudi Arabia thru 2010â€“2019&lt;/span&gt;. AIMS Public Health, 2020, 7, 679-696.	2.6	65
2	Î²â€“Actinâ€“dependent global chromatin organization and gene expression programs control cellular identity. FASEB Journal, 2018, 32, 1296-1314.	0.5	50
3	Nuclear myosin 1 contributes to a chromatin landscape compatible with RNA polymerase II transcription activation. BMC Biology, 2015, 13, 35.	3.8	48
4	Interferon-induced transmembrane protein-3 genetic variant rs12252 is associated with COVID-19 mortality. Genomics, 2021, 113, 1733-1741.	2.9	39
5	Targeted <i>SLC19A3</i> gene sequencing of 3000 Saudi newborn: a pilot study toward newborn screening. Annals of Clinical and Translational Neurology, 2019, 6, 2097-2103.	3.7	38
6	In Î²â€“actin knockouts, epigenetic reprogramming and rDNA transcription inactivation lead to growth and proliferation defects. FASEB Journal, 2016, 30, 2860-2873.	0.5	34
7	Biallelic variants in four genes underlying recessive osteogenesis imperfecta. European Journal of Medical Genetics, 2020, 63, 103954.	1.3	26
8	Mutated <i>RAP1GDS1</i> causes a new syndrome of dysmorphic feature, intellectual disability & speech delay. Annals of Clinical and Translational Neurology, 2020, 7, 956-964.	3.7	21
9	Nuclear Wiskottâ€“Aldrich syndrome protein co-regulates T cell factor 1-mediated transcription in T cells. Genome Medicine, 2017, 9, 91.	8.2	16
10	Biallelic variant in DACH1, encoding Dachshund Homolog 1, defines a novel candidate locus for recessive postaxial polydactyly type A. Genomics, 2021, 113, 2495-2502.	2.9	16
11	The effect of the VKORC1 promoter variant on warfarin responsiveness in the Saudi Warfarin Pharmacogenetic (SWAP) cohort. Scientific Reports, 2020, 10, 11613.	3.3	9
12	Pancytopenia, Recurrent Infection, Poor Wound Healing, Heterotopia of the Brain Probably Associated with A Candidate Novel de Novo CDC42 Gene Defect: Expanding the Molecular and Phenotypic Spectrum. Genes, 2021, 12, 294.	2.4	9
13	Blockade of p38 MAPK overcomes AML stem cell line KG1a resistance to 5-Fluorouridine and the impact on miRNA profiling. PLoS ONE, 2022, 17, e0267855.	2.5	7
14	A classification system for split-hand/ foot malformation (SHFM): A proposal based on 3 pedigrees with WNT10B mutations. European Journal of Medical Genetics, 2020, 63, 103738.	1.3	6
15	A novel interstitial deletion of chromosome 2q21.1â€“q23.3: Case report and literature review. Molecular Genetics & Genomic Medicine, 2020, 8, e1135.	1.2	6
16	Herbal melanin induces interleukinâ€“1Î² secretion and production by human THPâ€“1 monocytes via Tollâ€“like receptorâ€“2 and p38 MAPK activation. Experimental and Therapeutic Medicine, 2021, 22, 1081.	1.8	6
17	Identification of the TTC26 Splice Variant in a Novel Complex Ciliopathy Syndrome with Biliary, Renal, Neurological, and Skeletal Manifestations. Molecular Syndromology, 2021, 12, 133-140.	0.8	5
18	Blood pressureâ€“lowering activity of statins: a systematic literature review and meta-analysis of placebo-randomized controlled trials. European Journal of Clinical Pharmacology, 2020, 76, 1745-1754.	1.9	3

#	ARTICLE	IF	CITATIONS
19	Population pharmacokinetics of busulfan in Saudi pediatric patients undergoing hematopoietic stem cell transplantation. <i>International Journal of Clinical Pharmacy</i> , 2020, 42, 703-712.	2.1	3
20	Identification of Novel Mutations in Colorectal Cancer Patients Using AmpliSeq Comprehensive Cancer Panel. <i>Journal of Personalized Medicine</i> , 2021, 11, 535.	2.5	3
21	Identification of CSF3R Mutations in B-Lineage Acute Lymphoblastic Leukemia Using Comprehensive Cancer Panel and Next-Generation Sequencing. <i>Genes</i> , 2021, 12, 1326.	2.4	3